

CLAIMS

1. A method for isolating a full-length cDNA clone, the method comprising:
 - (a) determining a nucleotide sequence from the 5'-region of a cDNA clone contained in a cDNA library;
 - (b) determining the presence or absence of an initiation codon in the nucleotide sequence determined in (a) using an initiation codon prediction program; and
 - (c) selecting clones recognized as containing the initiation codon in (b).
2. The method of claim 1, wherein the cDNA library is constructed by a method for preparing a full length-enriched cDNA library.
3. The method of claim 1, wherein a cDNA library is constructed by a method comprising a step of modifying Cap of mRNA.
4. A method for constructing a full length cDNA library, the method comprising:
 - (a) determining a nucleotide sequence from the 5'-region of a cDNA clone contained in a cDNA library;
 - (b) determining the presence or absence of an initiation codon in the nucleotide sequence determined in (a) using an initiation codon prediction program;
 - (c) selecting clones recognized as containing the initiation codon in (b); and
 - (d) combining the clones selected in (c).
5. The method of claim 4, wherein the cDNA library is prepared by a method for constructing a full length-enriched cDNA library.
6. The method of claim 4, wherein the cDNA library is constructed by a method comprising a step of modifying Cap of mRNA.
7. A cDNA library obtainable by the method of claim 4.